SEQUENCE LISTING

<110> Kwon, Byoung S.

<120> MURINE 4-1BB GENE

<130> 740.009US1

<140> US 08/012,269
<141> 1993-02-01

<150> US 07/922,996
<151> 1992-07-30

<150> US 07/267,572
<151> 1988-11-07

<160> 13

<170> FastSEQ for Windows Version 4.0

<210> 1
<211> 2350
<212> DNA
<213> Mus musculus

T,0920

<220>
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<222> (1)...(2350)
<223> n = A,T,C or G

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acacacacac acacacacac acacacacgt ttatactacg tactgttatc ggtattctac 1560
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<210> 2
<211> 256
<212> PRT
<213> Mus musculus

<400> 2

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Pro Gly Thr Phe Cys Arg Lys Tyr Asn Pro Val Cys Lys Ser Cys Pro
        35                  40                  45
Pro Ser Thr Phe Ser Ser Ile Gly Gly Gln Pro Asn Cys Asn Ile Cys
    50                  55                  60
Arg Val Cys Ala Gly Tyr Phe Arg Phe Lys Lys Phe Cys Ser Ser Thr
65                  70                  75                  80
His Asn Ala Glu Cys Glu Cys Ile Glu Gly Phe His Cys Leu Gly Pro
                85                  90                  95
Gln Cys Thr Arg Cys Glu Lys Asp Cys Arg Pro Gly Gln Glu Leu Thr
            100                 105                 110
Lys Gln Gly Cys Lys Thr Cys Ser Leu Gly Thr Phe Asn Asp Gln Asn
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Gly Thr Gly Val Cys Arg Pro Trp Thr Asn Cys Ser Leu Asp Gly Arg
    130                 135                 140
Ser Val Leu Lys Thr Gly Thr Thr Glu Lys Asp Val Val Cys Gly Pro
145                 150                 155                 160
Pro Val Val Ser Phe Ser Pro Ser Thr Thr Ile Ser Val Thr Pro Glu
                165                 170                 175
Gly Gly Pro Gly Gly His Ser Leu Gln Val Leu Thr Leu Phe Leu Ala
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Leu Thr Ser Ala Leu Leu Leu Ala Leu Ile Phe Ile Thr Leu Leu Phe
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Ser Val Leu Lys Trp Ile Arg Lys Lys Phe Pro His Ile Phe Lys Gln
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<210> 3
<211> 24
<212> PRT
<213> Mus musculus

<400> 3
Cys Arg Val Cys Ala Gly Tyr Phe Arg Phe Lys Lys Phe Cys Ser Ser
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Thr His Asn Ala Glu Cys Glu Cys
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<210> 4
<211> 22
<212> PRT
<213> Drosophila

<400> 4
Cys Pro Val Cys Phe Asp Tyr Val Ile Leu Gln Cys Ser Ser Gly His
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Leu Val Cys Val Ser Cys
            20

<210> 5
<211> 26
<212> PRT
<213> Dictyostelium

<400> 5
Cys Pro Ile Cys Phe Glu Phe Ile Tyr Lys Lys Gln Ile Tyr Gln Cys
 1   5   10   15
Lys Ser Gly His His Ala Cys Lys Glu Cys
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<210> 6
<211> 6
<212> PRT
<213> Mus musculus

<220>
<221> SITE
<222> (1)...(6)
<223> Xaa = Any Amino Acid

<400> 6
Val Gln Asn Ser Xaa Asp
 1   5

<210> 7
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<223> An artificial peptide

<400> 7
Cys Arg Pro Gly Gln Glu Leu Thr Lys Ser Gly Tyr
 1   5   10

<210> 8
<211> 24
<212> PRT
<213> Artificial Sequence

<220>
<223> A conserved pattern

<221> SITE
<222> (1)...(24)
<223> Xaa = Any Amino Acid

<400> 8
Cys Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa
 1   5   10   15
Xaa His Xaa Xaa Xaa Cys Xaa Cys
            20

<210> 9
<211> 4
<212> PRT
<213> Mus musculus

<400> 9
Cys Arg Cys Pro
 1

<210> 10
<211> 4
<212> PRT
<213> Artificial Sequence

<220>
<223> A consensus sequence

<221> SITE
<222> (1)...(4)
<223> Xaa = Any Amino Acid

<400> 10
Cys Xaa Cys Pro
 1

<210> 11
<211> 25
<212> DNA
<213> Artificial Sequence

<220>
<223> A primer

<400> 11
acctcgaggt cctgtgcatg tgaca 25

<210> 12
<211> 25
<212> DNA
<213> Artificial Sequence

<220>
<223> A primer

<400> 12

atgaattctt actgcaggag tgccc 25

<210> 13
<211> 11
<212> PRT
<213> Mus musculus

<400> 13
Cys Arg Pro Gly Gln Glu Leu Thr Lys Gln Gly
1 5 10